

8: <u>P22392</u>

BLink, Domains, Links

Nucleoside diphosphate kinase B (NDK B) (NDP kinase B) (nm23-H2) (C-myc purine-binding transcription factor PUF) gi|127983|sp|P22392|NDKB HUMAN[127983]

9: <u>CAA53270</u>

BLink, Domains, Links

nm23H1g [Homo sapiens] gi|468542|emb|CAA53270.1|[468542]

10: CAA35621

BLink, Domains, Links

Nm23 protein [Homo sapiens] gi|35068|emb|CAA35621.1|[35068]

11: <u>NP_056542</u>

BLink, Domains, Links

tachykinin receptor 1 isoform short; neurokinin 1 receptor; Tachykinin receptor 1 (substance P receptor; neurokinin-1 receptor); NK-1 receptor; tachykinin 1 receptor (substance P receptor, neurokinin 1 receptor) [Homo sapiens]

gi|7669546|ref|NP_056542.1|[7669546]

12: NP 001049

BLink, Domains, Links

tachykinin receptor 1 isoform long; neurokinin 1 receptor; Tachykinin receptor 1 (substance P receptor; neurokinin-1 receptor); NK-1 receptor; tachykinin 1 receptor (substance P receptor, neurokinin 1 receptor) [Homo sapiens]

gi|4507343|ref|NP 001049.1|[4507343]

13: XP 488385

BLink, Links

similar to Regulator of G-protein signaling 3 (RGS3) [Mus musculus] gi|51772517|ref|XP 488385.1|[51772517]

14: XP 380045

BLink, Links

PREDICTED: similar to Nucleoside diphosphate kinase, mitochondrial precursor (NDP kinase, mitochondrial) (NDK) (nm23-H4) (Nucleoside diphosphate kinase D) (NDPKD) [Homo sapiens] gi|51492856|ref|XP 380045.2|[51492856]

15: XP 380047

BLink, Links

PREDICTED: similar to Nucleoside diphosphate kinase, mitochondrial precursor (NDP kinase, mitochondrial) (NDK) (nm23-H4) (Nucleoside diphosphate kinase D) (NDPKD) [Homo sapiens] gi|51492854|ref|XP 380047.2|[51492854]

16: NP 001001392

5)

BLink, Domains, Links

CD44 antigen isoform 5 precursor; cell surface glycoprotein CD44; Lutheran inhibitor, dominant; homing function and Indian blood group system; monoclonal antibody A3D8; antigen gp90 homing receptor; CDW44 antigen;

phagocytic glycoprotein I; extracellular matrix receptor-III; GP90 lymphocyte homing/adhesion receptor; heparan sulfate proteoglycan; cell adhesion molecule (CD44); hyaluronate receptor; Hermes antigen [Homo sapiens]

gi|48255943|ref|NP_001001392.1|[48255943]

17: NP 001001391

BLink, Domains, Links

CD44 antigen isoform 4 precursor; cell surface glycoprotein CD44; Lutheran inhibitor, dominant; homing function and Indian blood group system; monoclonal antibody A3D8; antigen gp90 homing receptor; CDW44 antigen; phagocytic glycoprotein I; extracellular matrix receptor-III; GP90 lymphocyte homing/adhesion receptor; heparan sulfate proteoglycan; cell adhesion molecule (CD44); hyaluronate receptor; Hermes antigen [Homo sapiens]

gi|48255941|ref|NP_001001391.1|[48255941]

18: NP 001001390

BLink, Domains, Links

CD44 antigen isoform 3 precursor; cell surface glycoprotein CD44; Lutheran inhibitor, dominant; homing function and Indian blood group system; monoclonal antibody A3D8; antigen gp90 homing receptor; CDW44 antigen; phagocytic glycoprotein I; extracellular matrix receptor-III; GP90 lymphocyte homing/adhesion receptor; heparan sulfate proteoglycan; cell adhesion molecule (CD44); hyaluronate receptor; Hermes antigen [Homo sapiens] gi[48255939]ref[NP_001001390.1][48255939]

19: NP 001001389

BLink, Domains, Links

CD44 antigen isoform 2 precursor; cell surface glycoprotein CD44; Lutheran inhibitor, dominant; homing function and Indian blood group system; monoclonal antibody A3D8; antigen gp90 homing receptor; CDW44 antigen; phagocytic glycoprotein I; extracellular matrix receptor-III; GP90 lymphocyte homing/adhesion receptor; heparan sulfate proteoglycan; cell adhesion molecule (CD44); hyaluronate receptor; Hermes antigen [Homo sapiens] gi[48255937]ref[NP_001001389.1][48255937]

20: NP 000601

BLink, Domains, Links

CD44 antigen isoform 1 precursor; cell surface glycoprotein CD44; Lutheran inhibitor, dominant; homing function and Indian blood group system; monoclonal antibody A3D8; antigen gp90 homing receptor; CDW44 antigen; phagocytic glycoprotein I; extracellular matrix receptor-III; GP90 lymphocyte homing/adhesion receptor; heparan sulfate proteoglycan; cell adhesion molecule (CD44); hyaluronate receptor; Hermes antigen [Homo sapiens]

gi|48255935|ref|NP 000601.3|[48255935]

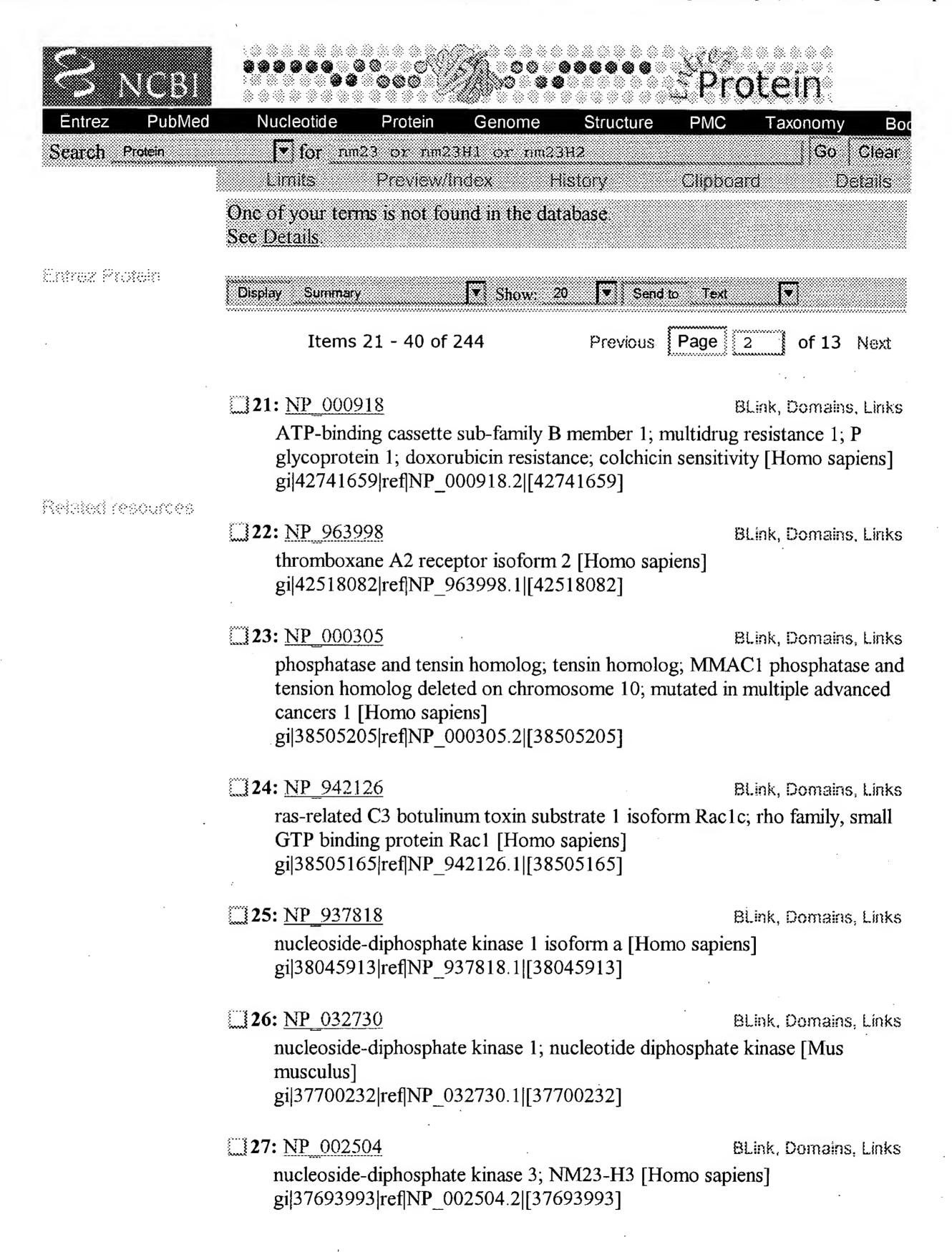
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BLink, Domains, Links

28: NP 872590 BLink, Domains, Links proliferating cell nuclear antigen; cyclin; DNA polymerase delta auxiliary protein [Homo sapiens] gi|33239451|ref|NP 872590.1|[33239451] **29:** NP 057700 BLink, Domains, Links NM23-H8; sperm-specific thioredoxin 2; thioredoxin domain-containing 3 (spermatozoa) [Homo sapiens] gi|31543836|ref|NP 057700.2|[31543836] **30:** NP 113751 BLink, Domains, Links nuclease sensitive element binding protein 1; Y box protein 1 [Rattus norvegicus] gi|31543347|ref|NP 113751.2|[31543347] **31:** NP 775482 BLink, Domains, Links PRUNEM1 [Mus musculus] gi|27597069|ref|NP 775482.1|[27597069] 32: NP 008845 BLink, Domains, Links RAR-related orphan receptor B; retinoic acid-binding receptor beta; nuclear receptor RZR-beta; RAR-related orphan receptor beta [Homo sapiens] gi|19743907|ref|NP 008845.2|[19743907] **33:** NP 599024 BLink, Domains, Links RAR-related orphan receptor A isoform d; retinoic acid receptor-related orphan receptor alpha; transcription factor RZR-alpha; ROR-alpha; RAR-related orphan receptor alpha [Homo sapiens] gi|19743905|ref|NP_599024.1|[19743905] **34:** NP 599023 BLink, Domains, Links RAR-related orphan receptor A isoform a; retinoic acid receptor-related orphan receptor alpha; transcription factor RZR-alpha; ROR-alpha; RAR-related orphan receptor alpha [Homo sapiens] gi|19743903|ref|NP_599023.1|[19743903] **35:** NP 599022 BLink, Domains, Links RAR-related orphan receptor A isoform b; retinoic acid receptor-related orphan receptor alpha; transcription factor RZR-alpha; ROR-alpha; RAR-related orphan receptor alpha [Homo sapiens] gi|19743901|ref|NP_599022.1|[19743901]

menin isoform 1; endocrine adenomatosis, multiple; Wermer syndrome;

Zollinger-Ellison syndrome, included; menin [Homo sapiens]

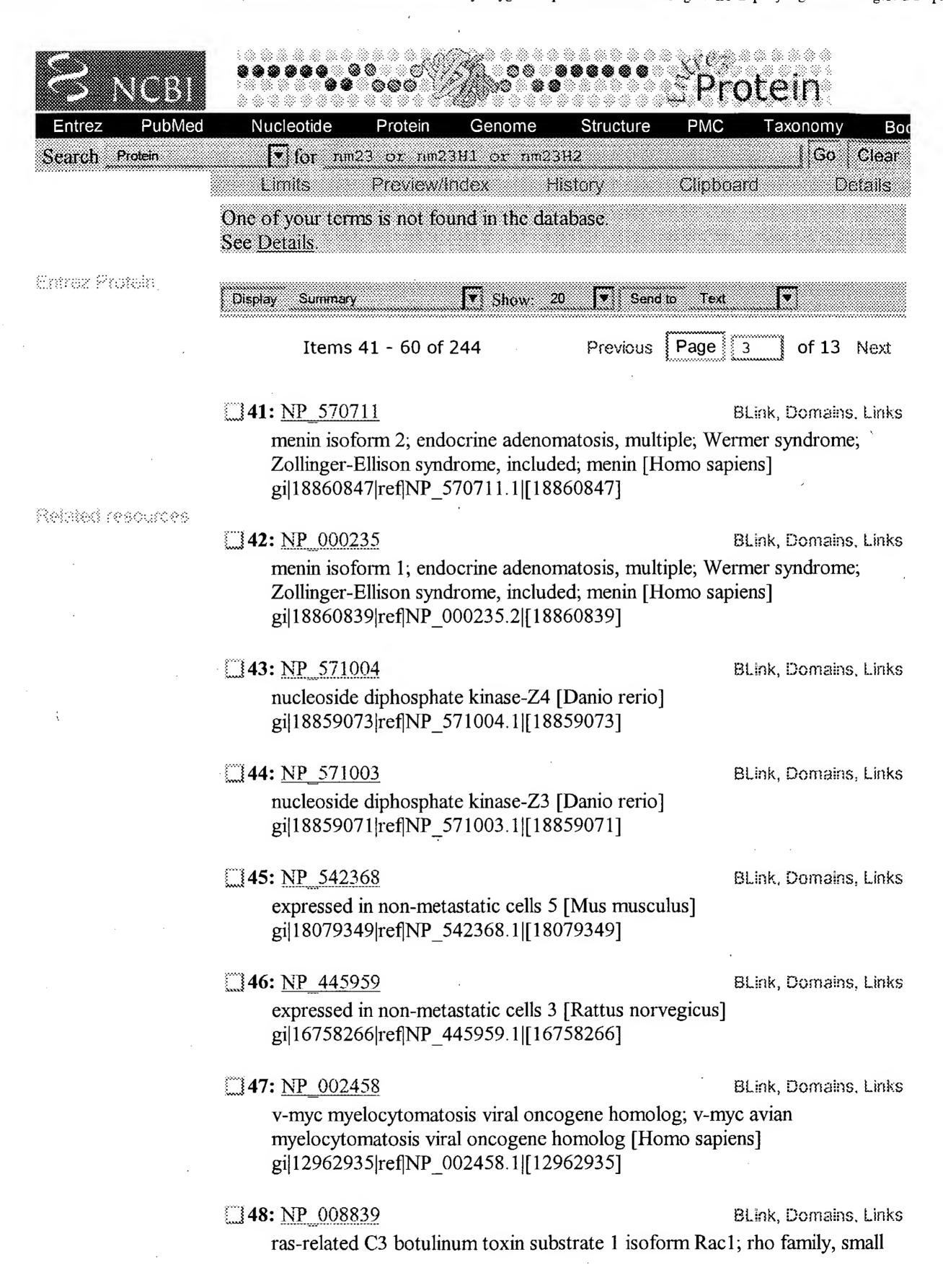
gi|18860857|ref|NP_570716.1|[18860857]

36: NP 570716

37: <u>NP_570715</u>	BLink, Domains, Links
menin isoform 1; endocrine adeno Zollinger-Ellison syndrome, includ gi 18860855 ref NP_570715.1 [18	
38: <u>NP_570714</u>	BLink, Domains, Links
menin isoform 1; endocrine adeno Zollinger-Ellison syndrome, includ gi 18860853 ref NP_570714.1 [18	
39: <u>NP 570713</u>	BLink, Domains, Links
menin isoform 1; endocrine adeno Zollinger-Ellison syndrome, includ gi 18860851 ref NP_570713.1 [18	
40: NP 570712	BLink, Domains, Links
menin isoform 1; endocrine adeno Zollinger-Ellison syndrome, includ gi 18860849 ref NP_570712.1 [18	1 3
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GTP binding protein Rac1 [Homo sapiens] gi|9845511|ref|NP_008839.2|[9845511]

149: NP 061485

BLink, Domains, Links

ras-related C3 botulinum toxin substrate 1 isoform Rac1b; rho family, small GTP binding protein Rac1 [Homo sapiens] gi[9845509|ref|NP 061485.1|[9845509]

50: NP_062705

BLink, Domains, Links

nucleoside diphosphate kinase 4; nucleoside diphosphate kinase [Mus musculus]

gi|9790123|ref|NP_062705.1|[9790123]

51: NP 062704

BLink, Domains, Links

nucleoside diphosphate kinase DR-nm23 [Mus musculus] gi|9790121|ref|NP 062704.1|[9790121]

52: <u>NP_061227</u>

BLink, Domains, Links

expressed in non-metastatic cells 6, protein; nucleoside diphosphate kinase type 6; nm23/nucleoside diphosphate kinase 6; expressed in non-metastatic cells 6, protein (nucleoside diphosphate kinase) [Mus musculus] gi|9055290|ref|NP 061227.1|[9055290]

53: NP 000537

BLink, Domains, Links

tumor protein p53 [Homo sapiens] gi|8400738|ref|NP 000537.2|[8400738]

54: NP 038599

BLink, Domains, Links

kinase suppressor of ras [Mus musculus] gi|7305215|ref|NP 038599.1|[7305215]

55: NP 032731

BLink, Domains, Links

nucleoside-diphosphate kinase 2; nucleoside diphosphate kinase B [Mus musculus]

gi|6679078|ref|NP_032731.1|[6679078]

56: NP_006370

BLink, Domains, Links

semaphorin 3C; semaphorin E [Homo sapiens] gi|5454048|ref|NP 006370.1|[5454048]

57: NP 005000

BLink, Domains, Links

nucleoside-diphosphate kinase 4 [Homo sapiens] gi|4826862|ref|NP 005000.1|[4826862]

58: NP_004439

Bl.ink, Domains, Links

v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog; Avian erythroblastic

leukemia viral (v-erb-b2) oncogene homolog 2; v-erb-b2 avian erythroblastic

leukemia viral oncogene homolog 2 (neuro/glioblastoma derived oncogene

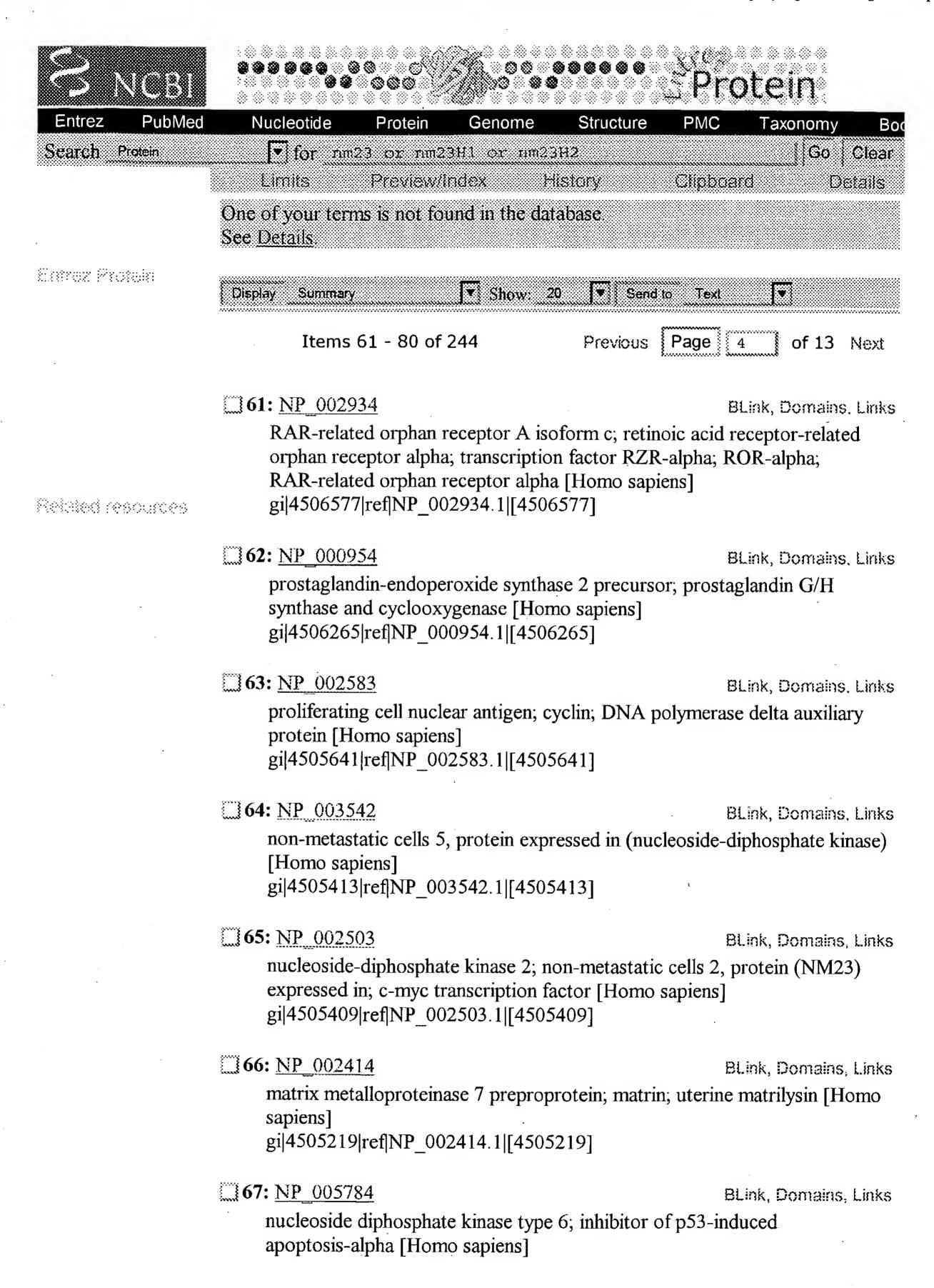
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gi|5031951|ref|NP 005784.1|[5031951]

68: NP 067045

BLink, Domains, Links

TcD37 homolog; prune [Homo sapiens] gi|24308263|ref|NP_067045.1|[24308263]

69: NP_006401

BLink, Domains, Links

HIV-1 Tat interactive protein 2, 30kDa; Tat-interacting protein (30kD); HIV-1 Tat interactive protein 2, 30 kDa; HIV-1 Tat interactive protein 2, 30 kD [Homo sapiens] gi|20127503|ref|NP_006401.2|[20127503]

70: NP 002408

Blink, Domains, Links

antigen identified by monoclonal antibody Ki-67; Proliferation-related Ki-67 antigen [Homo sapiens] gi|19923217|ref|NP_002408.2|[19923217]

71: NP_112316

BLink, Domains, Links

matrix metalloproteinase 2 (72 KDa type IV collagenase); gelatinase A [Rattus norvegicus] gi|13591991|ref|NP_112316.1|[13591991]

72: NP 006656

BLink, Domains, Links

heparanase; heparanase-1 [Homo sapiens] gi|5729873|ref|NP_006656.1|[5729873]

73: XP 377976

BLink, Links

PREDICTED: similar to Nucleoside diphosphate kinase, mitochondrial precursor (NDP kinase, mitochondrial) (NDK) (nm23-H4) (Nucleoside diphosphate kinase D) (NDPKD) [Homo sapiens] gi|51466625|ref|XP_377976.2|[51466625]

74: EAL24027

BLink, Links

similar to Nucleoside diphosphate kinase, mitochondrial precursor (NDP kinase, mitochondrial) (NDK) (nm23-M4) (Nucleoside diphosphate kinase D) [Homo sapiens] gi|51094781|gb|EAL24027.1|[51094781]

75: XP 424348

BLink, Links

PREDICTED: similar to Nucleoside diphosphate kinase homolog 5 (NDK-H 5) (NDP kinase homolog 5) (nm23-H5) (Testis-specific nm23 homolog) (Inhibitor of p53-induced apoptosis-beta) (IPIA-beta) [Gallus gallus] gi|50805547|ref|XP_424348.1|[50805547]

76: <u>XP 417888</u>

BLink, Links

PREDICTED: similar to nm23-phosphorylated unknown substrate; SH3 domain-containing 70 kDa protein [Gallus gallus]

PREDICTED: similar to Nucleoside diphosphate kinase homolog 5 (NDK-H 5) (NDP kinase homolog 5) (nm23-H5) (Testis-specific nm23 homolog) (Inhibitor of p53-induced apoptosis-beta) (IPIA-beta) [Gallus gallus] gi|50755299|ref|XP_414687.1|[50755299]

PREDICTED: similar to NM23-H8; sperm-specific thioredoxin 2; thioredoxin domain-containing 3 (spermatozoa) [Gallus gallus] gi|50733086|ref|XP_426021.1|[50733086]

gi|50760071|ref|XP 417888.1|[50760071]

PREDICTED: similar to Nucleoside diphosphate kinase 7 (NDK 7) (NDP kinase 7) (nm23-H7) [Gallus gallus] gi|50729630|ref|XP_416591.1|[50729630]

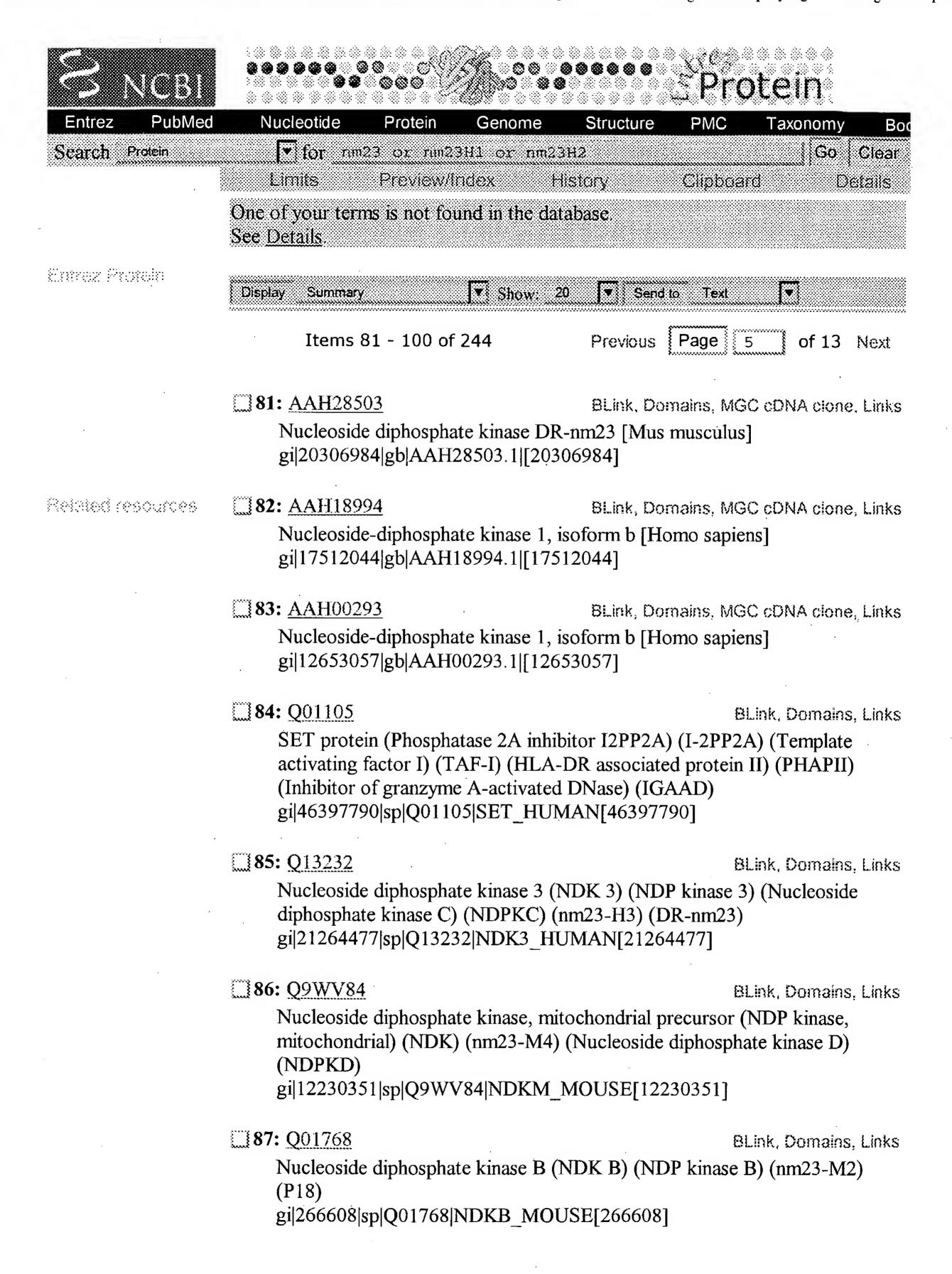
80: <u>AAH36816</u>
NM23-H8 [Homo sapiens]
gi|22477642|gb|AAH36816.1|[22477642]

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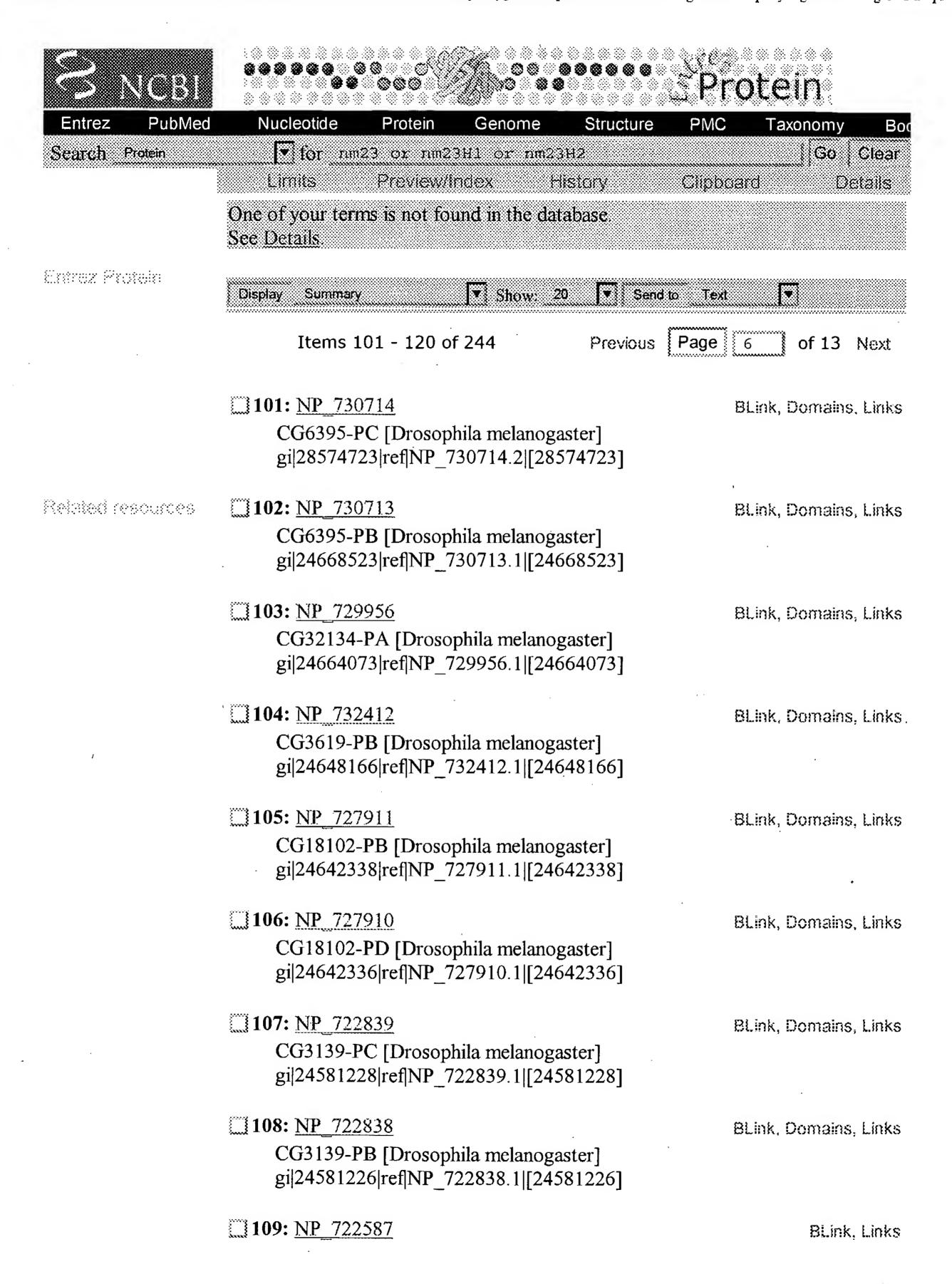


88: <u>P15532</u> BLink, Domains, Links Nucleoside diphosphate kinase A (NDK A) (NDP kinase A) (Tumor metastatic process-associated protein) (Metastasis inhibition factor NM23) (NDPK-A) (nm23-M1) gi|127982|sp|P15532|NDKA MOUSE[127982] **89:** XP 396235 BLink, Domains, Links similar to Nucleoside diphosphate kinase 7 (NDK 7) (NDP kinase 7) (nm23-R7) [Apis mellifera] gi|48109475|ref|XP 396235.1|[48109475] **90:** XP 396140 BLink, Domains, Links similar to Nucleoside diphosphate kinase 6 (NDK 6) (NDP kinase 6) (nm23-H6) (Inhibitor of p53-induced apoptosis-alpha) (IPIA-alpha) [Apis mellifera] gi|48106672|ref|XP_396140.1|[48106672] **91:** XP 394702 BLink, Domains, Links similar to Nucleoside diphosphate kinase homolog 5 (NDK-H 5) (NDP kinase homolog 5) (nm23-H5) (Testis-specific nm23 homolog) (Inhibitor of p53-induced apoptosis-beta) (IPIA-beta) [Apis mellifera] gi|48096513|ref|XP 394702.1|[48096513] **92:** XP 394464 BLink, Domains, Links similar to multiple endocrine neoplasia type 1 [Apis mellifera] gi|48095533|ref|XP 394464.1|[48095533] **93:** Q9WV85 BLink, Domains, Links Nucleoside diphosphate kinase 3 (NDK 3) (NDP kinase 3) (Nucleoside diphosphate kinase C) (NDPKC) (nm23-M3) (DR-nm23) gi|48429268|sp|Q9WV85|NDK3 MOUSE[48429268] **94:** AAH02664 **BLink**, Domains, Links MEN1 protein [Homo sapiens] gi|38197214|gb|AAH02664.2|[38197214] **95:** <u>AAH33449</u> BLink, Domains, MGC cDNA clone, Links Rgs3 protein [Mus musculus] gi|23270966|gb|AAH33449.1|[23270966] **96:** NP 996468 BLink, Domains, Links CG18102-PA [Drosophila melanogaster] gi|45555521|ref|NP 996468.1|[45555521] **97:** NP 996467 BLink, Domains, Links CG18102-PE [Drosophila melanogaster] gi|45555505|ref|NP 996467.1|[45555505]

98: NP 996466	BLink, Domains, Links
. CG18102-PF [Drosophila melanogas	ter]
gi 45555485 ref NP_996466.1 [45555	5485]
99: NP_996465	BLink, Domains, Links
CG18102-PG [Drosophila melanogas	ster]
gi 45555473 ref NP_996465.1 [45555	5473]
100: NP 995619	BLink, Domains, Links
CG3139-PD [Drosophila melanogast	er]
gi 45552193 ref NP_995619.1 [45552	2193]
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CG2671-PF [Drosophila melanogaster] gi|24580507|ref|NP_722587.1|[24580507]

110: NP 722586

BLink, Links

CG2671-PE [Drosophila melanogaster] gi|24580505|ref|NP_722586.1|[24580505]

111: NP 722585

Bl.ink, Links

CG2671-PD [Drosophila melanogaster] gi|24580503|ref|NP_722585.1|[24580503]

112: NP 722584

BLink, Links

CG2671-PA [Drosophila melanogaster] gi|24464586|ref|NP | 722584.1|[24464586]

113: NP 722583

BLink, Links

CG2671-PB [Drosophila melanogaster] gi|24464584|ref|NP_722583.1|[24464584]

114: NP_523485

BLink, Domains, Links

CG7234-PI [Drosophila melanogaster] gi|45549153|ref|NP_523485.3|[45549153]

115: NP 476761

BLink, Domains, Links

CG2210-PA [Drosophila melanogaster] gi|45549037|ref|NP_476761.2|[45549037]

116: NP 523872

BLink, Domains, Links

CG6883-PA [Drosophila melanogaster] gi|24654763|ref|NP_523872.2|[24654763]

117: NP 523371

BLink, Domains, Links

CG9907-PA [Drosophila melanogaster] gi|24642537|ref|NP_523371.2|[24642537]

118: NP 524853

BLink, Domains, Links

CG18102-PC [Drosophila melanogaster] gi|24642340|ref|NP_524853.2|[24642340]

119: NP 511119

BLink, Domains, Links

CG1594-PA [Drosophila melanogaster] gi|24641273|ref|NP 511119.2|[24641273]

120: NP 476859

BLink, Domains, Links

CG3936-PA [Drosophila melanogaster] gi|24639454|ref|NP_476859.2|[24639454]

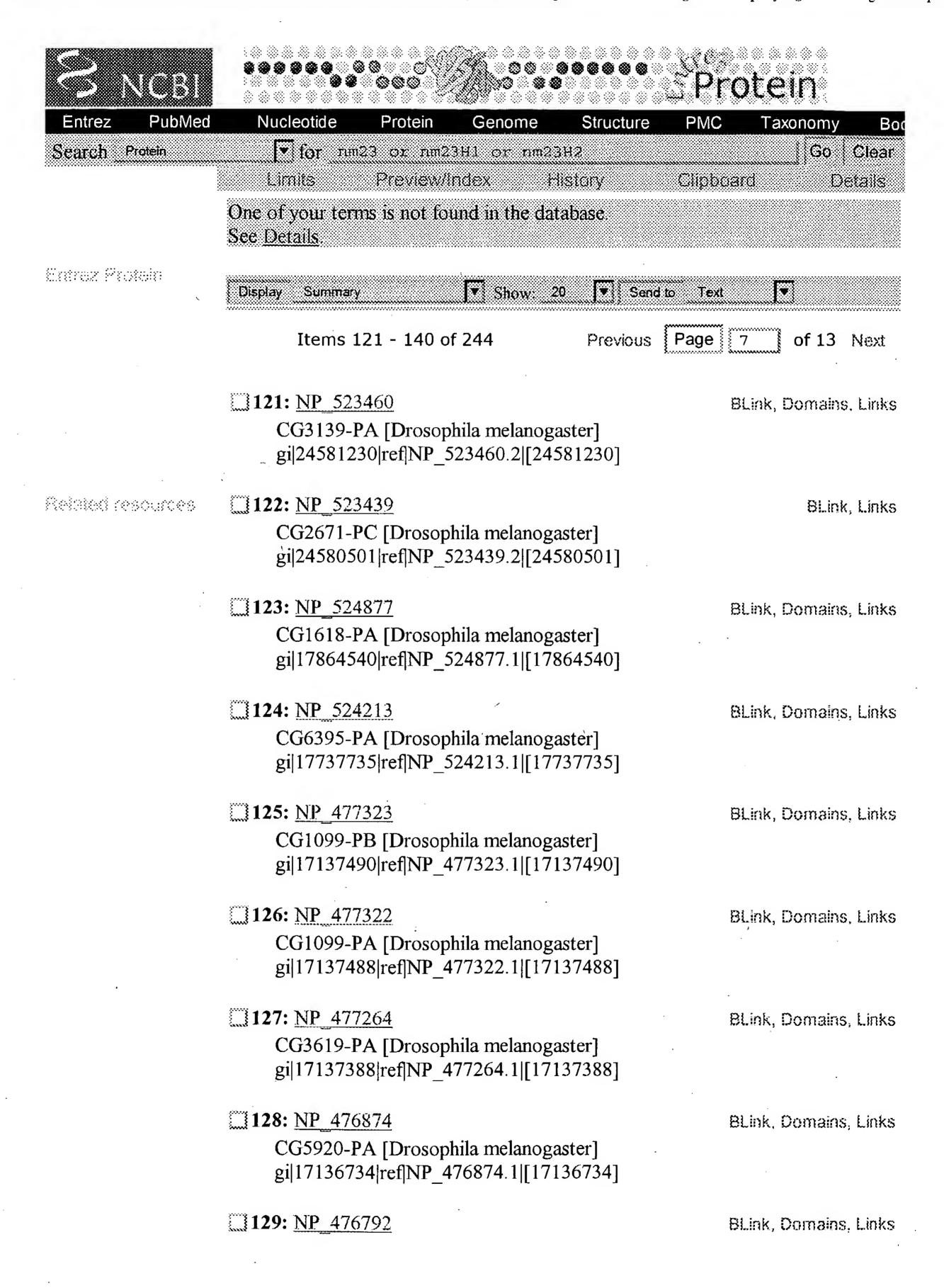
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CG3757-PA [Drosophila melanogaster] gi|17136600|ref|NP 476792.1|[17136600] **130:** NP 476759 BLink, Domains, Links CG10079-PB [Drosophila melanogaster] gi|17136536|ref|NP 476759.1|[17136536] **131:** NP 476758 BLink, Domains, Links CG10079-PA [Drosophila melanogaster] gi|17136534|ref|NP 476758.1|[17136534] **132:** NP 476684 BLink, Domains, Links CG3461-PA [Drosophila melanogaster] gi|17136406|ref|NP_476684.1|[17136406] **133:** Q05982 BLink, Domains, Links Nucleoside diphosphate kinase A (NDK A) (NDP kinase A) (Tumor metastatic process-associated protein) (Metastasis inhibition factor NM23) gi|462690|sp|Q05982|NDKA RAT[462690] **134:** <u>BAC39392</u> BLink, Domains, Links unnamed protein product [Mus musculus] gi|26351511|dbj|BAC39392.1|[26351511] **135:** <u>BAB30896</u> BLink, Domains, Links unnamed protein product [Mus musculus] gi|12857112|dbj|BAB30896.1|[12857112] **136:** BAB27708 BLink, Links unnamed protein product [Mus musculus] gi|12847786|dbi|BAB27708.1|[12847786] **137:** NP 173184 BLink, Domains, Links nucleoside diphosphate kinase family protein [Arabidopsis thaliana] gi|42562123|ref|NP_173184.2|[42562123] **138:** 075414 BLink; Domains, Links Nucleoside diphosphate kinase 6 (NDK 6) (NDP kinase 6) (nm23-H6) (Inhibitor of p53-induced apoptosis-alpha) (IPIA-alpha) gi|12232627|sp|O75414|NDK6_HUMAN[12232627] **139:** NP 508832 BLink, Domains, Links nm23-phosphorylated substrate (XF511) [Caenorhabditis elegans] gi|17569663|ref|NP 508832.1|[17569663] **140:** <u>AAQ77892</u> BLink, Links

Sequence 5 from patent US 6329198

gi|34599178|gb|AAQ77892.1||pat|US|6329198|5[34599178]

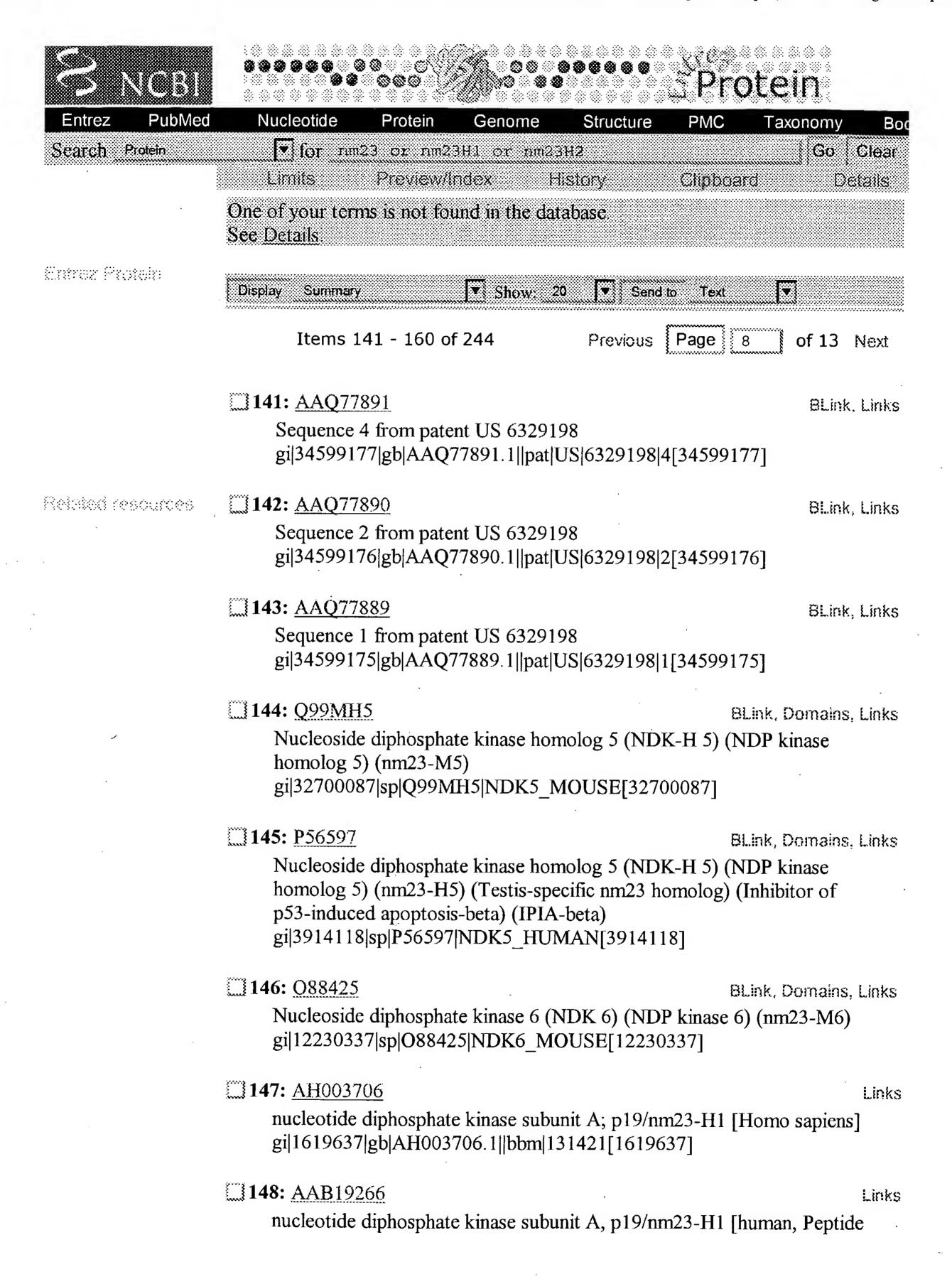
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BLink, Domains, Links

Partial, 6 aa, segment 3 of 3] gi|232476|gb|AAB19266.1||bbs|38818[232476] **149:** <u>AAB19265</u> Links No definition line found gi|232475|gb|AAB19265.1||bbs|38817[232475] **150:** AAB19264 Links nucleotide diphosphate kinase subunit A, p19/nm23-H1 [human, Peptide Partial, 10 aa, segment 2 of 3] gi|232474|gb|AAB19264.1||bbs|38815[232474] **151:** <u>AAB19263</u> Links No definition line found gi|232473|gb|AAB19263.1||bbs|38813[232473] **152:** AAB19262 Links nucleotide diphosphate kinase subunit A, p19/nm23-H1 [human, Peptide Partial, 12 aa, segment 1 of 3] gi|232472|gb|AAB19262.1||bbs|38811[232472] **153:** AAO85436 BLink, Domains, Links NM23-H1 [Homo sapiens] gi|29468184|gb|AAO85436.1|AF487339_1[29468184] **154:** <u>AAC95290</u> BLink, Domains, Links PRUNE-like protein [Homo sapiens] gi|4007408|gb|AAC95290.1|[4007408] **155:** <u>1JXVF</u> BLink, Domains, Links Chain F, Crystal Structure Of Human Nucleoside Diphosphate Kinase A gi|20663972|pdb|1JXV|F[20663972] **156:** 1JXVE BLink, Domains, Links Chain E, Crystal Structure Of Human Nucleoside Diphosphate Kinase A gi|20663971|pdb|1JXV|E[20663971] **157:** <u>1JXVD</u> BLink, Domains, Links Chain D, Crystal Structure Of Human Nucleoside Diphosphate Kinase A gi|20663970|pdb|1JXV|D[20663970] **158:** 1JXVC BLink, Domains, Links Chain C, Crystal Structure Of Human Nucleoside Diphosphate Kinase A gi|20663969|pdb|1JXV|C[20663969]

Chain B, Crystal Structure Of Human Nucleoside Diphosphate Kinase A

159: <u>1JXVB</u>

gi|20663968|pdb|1JXV|B[20663968]

[[] 160: 1JXVA

BLink, Domains, Links

Chain A, Crystal Structure Of Human Nucleoside Diphosphate Kinase A gi|20663967|pdb|1JXV|A[20663967]

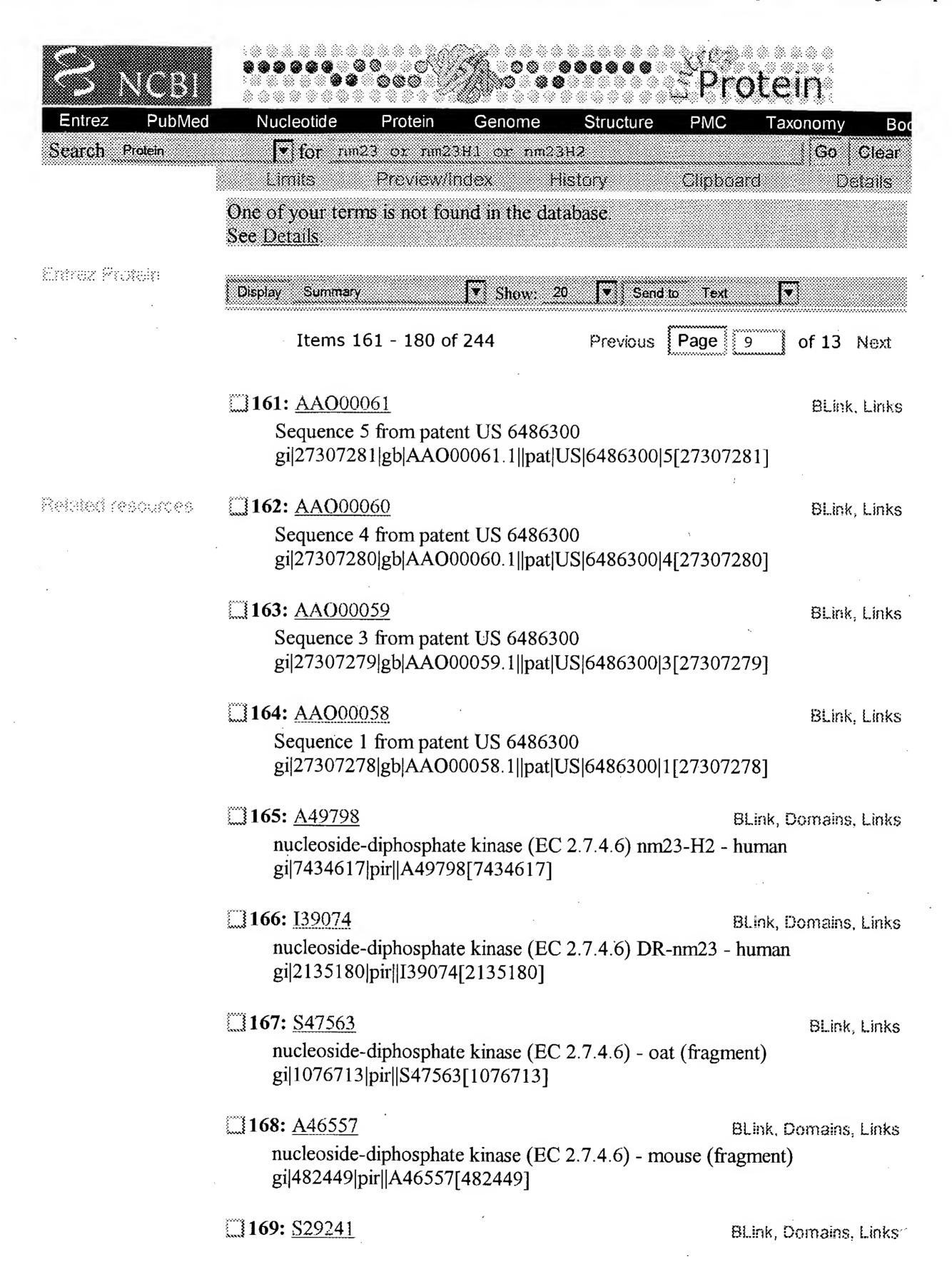
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-Sursiyaalik Joli



nucleoside-diphosphate kinase (EC 2.7.4.6) B - mouse gi|478765|pir||S29241[478765]

170: <u>A33386</u> BLink, Domains, Links nucleoside-diphosphate kinase (EC 2.7.4.6) nm23-H1g - human gi|88268|pir||A33386[88268] **171:** AAG02202 BLink, Domains, Links nucleoside diphosphate kinase D [Mus musculus] gi|9931518|gb|AAG02202.1|AF288692_1[9931518] **172:** AAG02201 BLink, Domains, Links nucleoside diphosphate kinase C [Mus musculus] gi|9931516|gb|AAG02201.1|AF288691 1[9931516] **173:** AAG02200 BLink, Domains, Links nucleoside diphosphate kinase D [Mus musculus] gi|9931514|gb|AAG02200.1|AF288690 1[9931514] **174:** AAG02199 BLink, Domains, Links nucleoside diphosphate kinase C [Mus musculus] gi|9931512|gb|AAG02199.1|AF288689_1[9931512] **175:** AAN23827 BLink, Links Sequence 3 from patent US 6423836 gi|23325147|gb|AAN23827.1||pat|US|6423836|3[23325147] **176:** <u>AAN23826</u> **BLink**, Links Sequence 1 from patent US 6423836 gi|23325146|gb|AAN23826.1||pat|US|6423836|1[23325146] **177:** <u>088426</u> BLink, Domains, Links Nucleoside diphosphate kinase 6 (NDK 6) (NDP kinase 6) (nm23-R6) gi|12230330|sp|O88426|NDK6_RAT[12230330] **178:** <u>AAL16953</u> BLink, Domains, Links nm23-phosphorylated unknown substrate [Homo sapiens] gi|16304176|gb|AAL16953.1|AF425252 1[16304176] **179:** AAF74448 BLink, Domains, Links nucleoside diphosphate kinase NDPK-Z6 [Danio rerio] gi|8308035|gb|AAF74448.1|AF241153 1[8308035] **180:** AAF60971 BLink, Domains, Links nuclease diphosphate kinase B [Danio rerio]

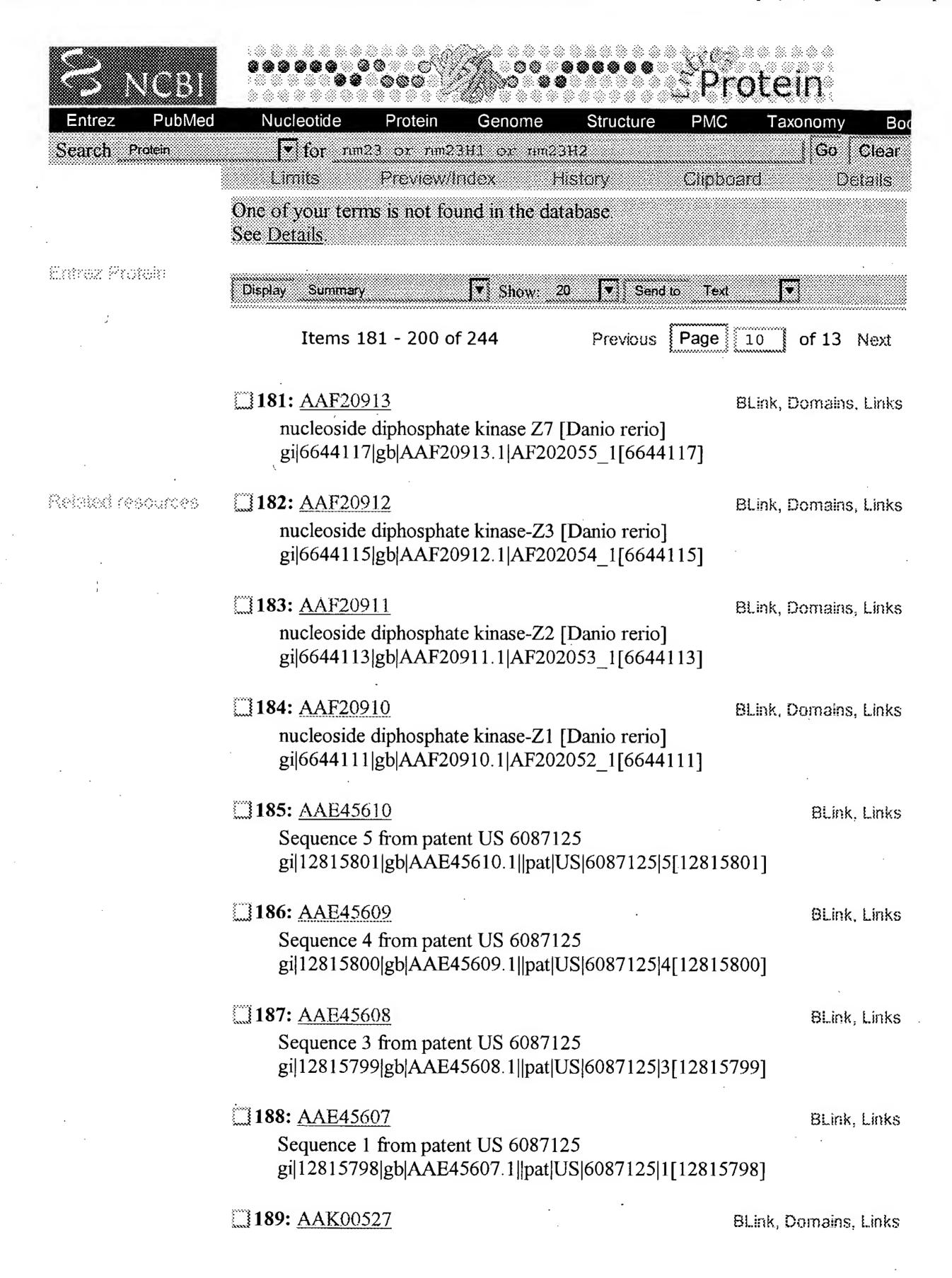
gi|7339840|gb|AAF60971.1|AF201764 1[7339840]

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nucleoside diphosphate kinase A [Cavia porcellus] gi|12700713|gb|AAK00527.1|[12700713]

190: <u>AAG54075</u> BLink, Domains, Links nucleoside diphosphate kinase DR-nm23 [Rattus norvegicus] gi|12621064|gb|AAG54075.1|[12621064] **191:** AAG14350 BLink, Domains, Links putative oncoprotein nm23 [Ictalurus punctatus] gi|10180968|gb|AAG14350.1|AF283993_1[10180968] **192:** 1EHWB BLink, Domains, Links Chain B, Human Nucleoside Diphosphate Kinase 4 gi|8569509|pdb|1EHW|B[8569509] **193:** 1EHWA BLink, Domains, Links Chain A, Human Nucleoside Diphosphate Kinase 4 gi|8569508|pdb|1EHW|A[8569508] **194:** <u>P70011</u> BLink, Domains, Links NUCLEOSIDE DIPHOSPHATE KINASE A2 (NDK A2) (NDP KINASE A2) (NM23/NUCLEOSIDE DIPHOSPHATE KINASE A2) gi|6225752|sp|P70011|NDK2_XENLA[6225752] **195:** <u>P70010</u> BLink, Domains, Links NUCLEOSIDE DIPHOSPHATE KINASE A1 (NDK A1) (NDP KINASE A1) (NM23/NUCLEOSIDE DIPHOSPHATE KINASE A1) gi|6225751|sp|P70010|NDK1 XENLA[6225751] **___196:** <u>AAF64467</u> BLink, Domains, Links type 6 nucleoside diphosphate kinase [Drosophila melanogaster] gi|7595825|gb|AAF64467.1|AF241151_1[7595825] **197:** AAF20909 BLink, Domains, Links NM23-H8 [Homo sapiens] gi|7580490|gb|AAF20909.2|AF202051_1[7580490] **198:** <u>AAD34622</u> BLink, Domains, Links nm23-H7 [Homo sapiens] gi|4960169|gb|AAD34622.1|AF153191 1[4960169] **199:** AAF20908 BLink, Domains, Links nmdyn-D7 [Drosophila melanogaster] gi|6644107|gb|AAF20908.1|AF202050 1[6644107] **200:** <u>AAF20907</u> BLink, Domains, Links NM23-R7 [Rattus norvegicus]

gi|6644105|gb|AAF20907.1|AF202049 1[6644105]

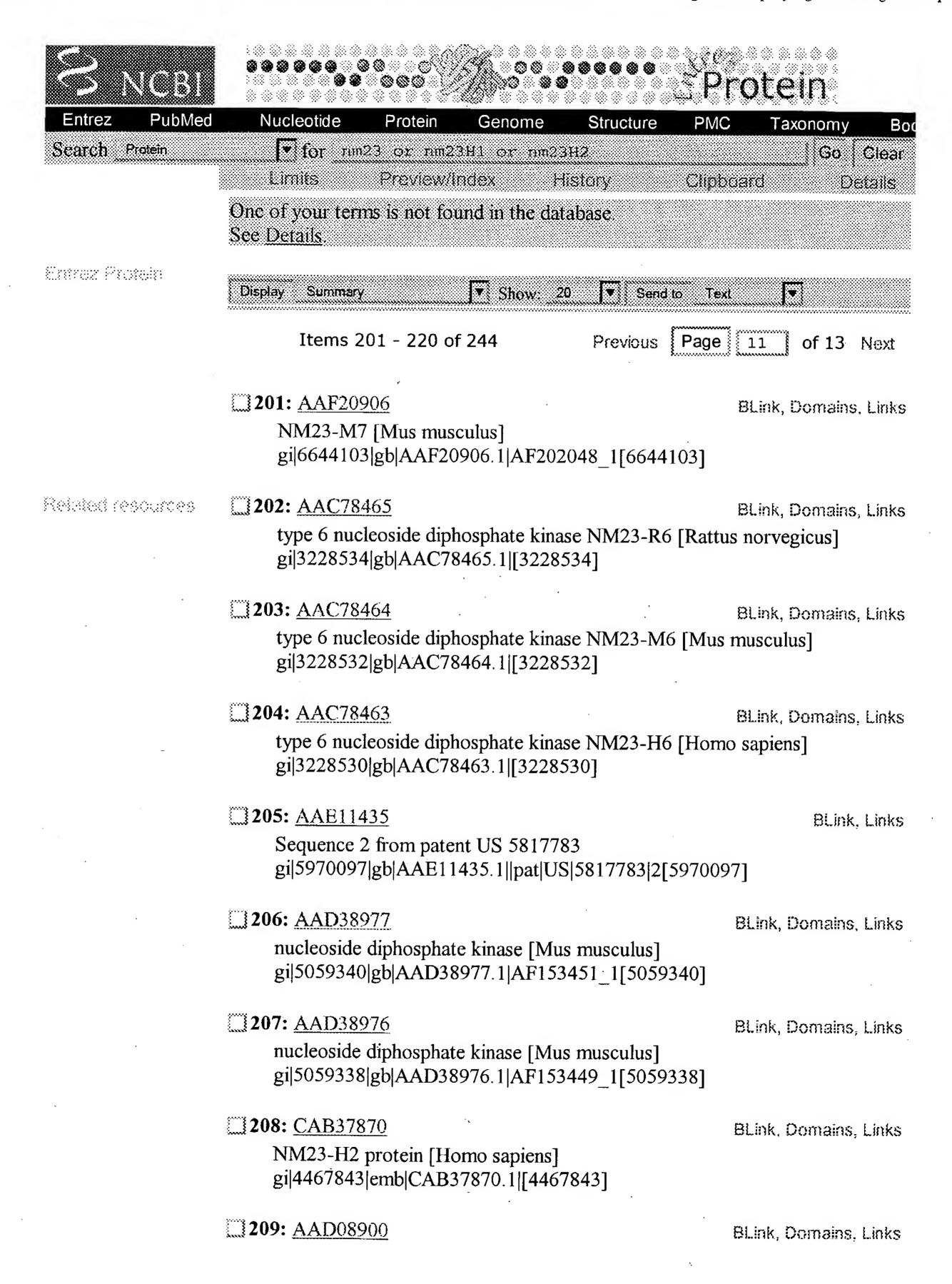
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nucleoside diphosphate kinase; NDP kinase [Scyliorhinus torazame] gi|4176739|gb|AAD08900.1|[4176739]

210: AAC64358

BLink, Domains, Links

nm23-H5 [Homo sapiens] gi|3220239|gb|AAC64358.1|[3220239]

211: AAC44154

BLink, Domains, Links

Ndk gi|1353658|gb|AAC44154.1|[1353658]

212: CAA75226

BLink, Domains, Links

nucleoside-diphosphate kinase [Homo sapiens] gi|3559927|emb|CAA75226.1|[3559927]

213: 1NUEF

BLink, Domains, Links

Chain F, Nucleoside Triphosphate, Nucleoside Diphosphate Mol_id: 1; Molecule: Nucleoside Diphosphate Kinase; Chain: A, B, C, D, E, F; Ec: 2.7.4.6 gi|1421614|pdb|1NUE|F[1421614]

214: 1NUEE

BLink, Domains, Links

Chain E, Nucleoside Triphosphate, Nucleoside Diphosphate Mol_id: 1; Molecule: Nucleoside Diphosphate Kinase; Chain: A, B, C, D, E, F; Ec: 2.7.4.6 gi[1421613]pdb[1NUE]E[1421613]

215: 1NUED

BLink, Domains, Links

Chain D, Nucleoside Triphosphate, Nucleoside Diphosphate Mol_id: 1; Molecule: Nucleoside Diphosphate Kinase; Chain: A, B, C, D, E, F; Ec: 2.7.4.6 gi|1421612|pdb|1NUE|D[1421612]

216: <u>1NUEC</u>

BLink, Domains, Links

Chain C, Nucleoside Triphosphate, Nucleoside Diphosphate Mol_id: 1; Molecule: Nucleoside Diphosphate Kinase; Chain: A, B, C, D, E, F; Ec: 2.7.4.6 gi[1421611]pdb[1NUE]C[1421611]

217: <u>1NUEB</u>

BLink, Domains, Links

Chain B, Nucleoside Triphosphate, Nucleoside Diphosphate Mol_id: 1; Molecule: Nucleoside Diphosphate Kinase; Chain: A, B, C, D, E, F; Ec: 2.7.4.6 gi[1421610]pdb[1NUE]B[1421610]

218: <u>INUEA</u>

BLink, Domains, Links

Chain A, Nucleoside Triphosphate, Nucleoside Diphosphate Mol id: 1;

Molecule: Nucleoside Diphosphate Kinase; Chain: A, B, C, D, E, F; Ec: 2.7.4.6 gi[1421609]pdb[1NUE]A[1421609]

219: <u>1NSKO</u>

BLink, Domains, Links

Chain O, Mol_id: 1; Molecule: Nucleoside Diphosphate Kinase; Chain: R, L, T, U, N, O; Ec: 2.7.4.6; Engineered: Yes gi|1311292|pdb|1NSK|0[1311292]

220: <u>1NSKN</u>

BLink, Domains, Links

Chain N, Mol_id: 1; Molecule: Nucleoside Diphosphate Kinase; Chain: R, L, T, U, N, O; Ec: 2.7.4.6; Engineered: Yes gi[1311291]pdb[1NSK]N[1311291]

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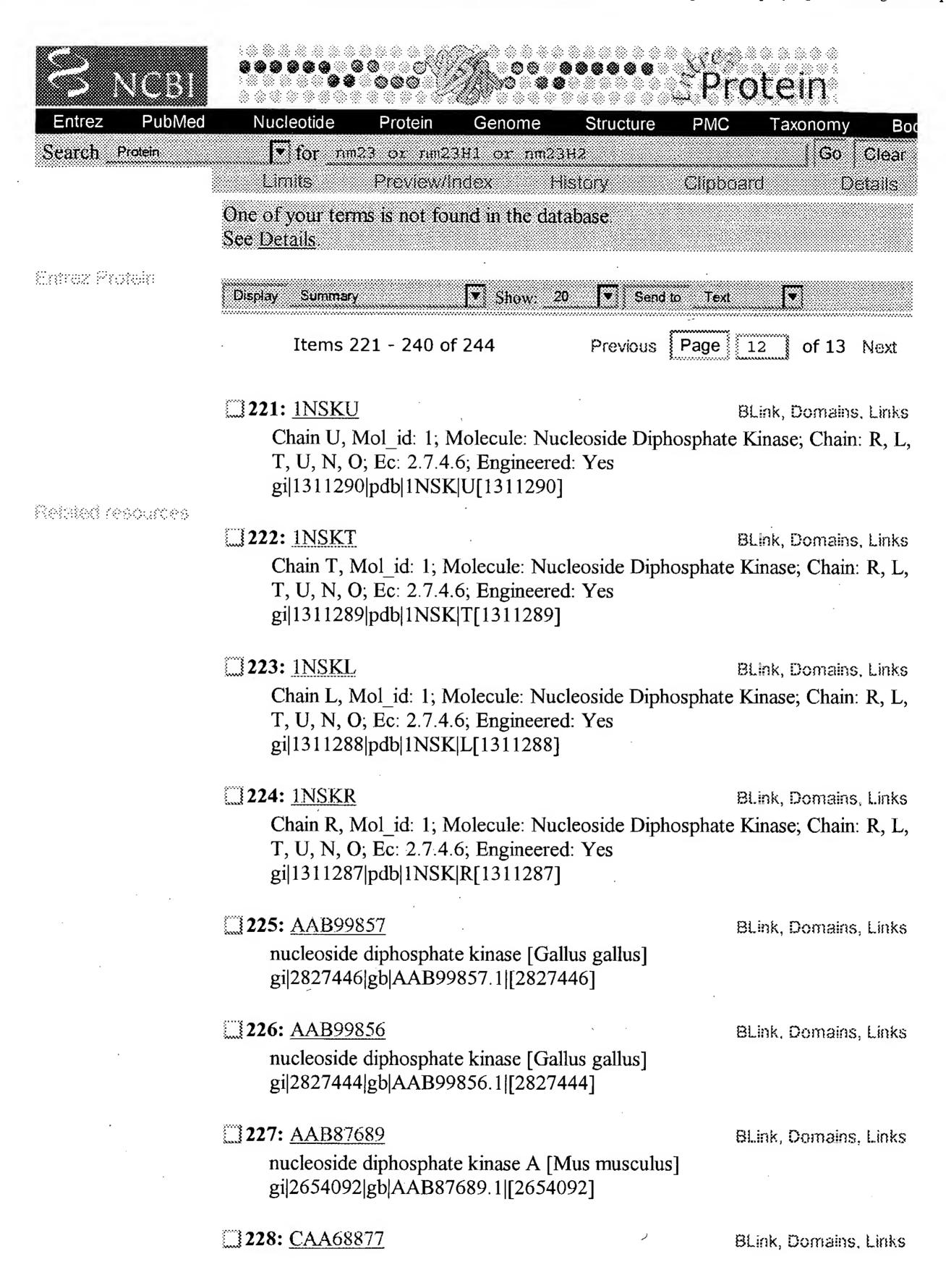
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nucleoside-diphosphate kinase [Homo sapiens] gi|1945762|emb|CAA68877.1|[1945762]

229: <u>AAB42080</u>

BLink, Domains, Links

nucleoside diphosphate kinase A long form [Mus musculus] gi|1816594|gb|AAB42080.1|[1816594]

230: <u>CAA66475</u>

BLink, Domains, Links

NM23/nucleoside diphosphate kinase [Xenopus laevis] gi|1655710|emb|CAA66475.1|[1655710]

231: CAA66476

BLink, Domains, Links

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232: CAA66474

BLink, Domains, Links

NM23/nucleoside diphosphate kinase [Xenopus laevis] gi|1655706|emb|CAA66474.1|[1655706]

233: CAA66473

BLink, Domains, Links

NM23/nucleoside diphosphate kinase [Xenopus laevis] gi|1655704|emb|CAA66473.1|[1655704]

234: CAA48275

BLink, Domains, Links

nucleoside diphosphate kinase B [Mus musculus] gi|53354|emb|CAA48275.1|[53354]

235: AAB31385

BLink, Links

nucleoside diphosphate kinase, NDPK=Nm23 protein homolog {N-terminal} {EC 2.7.4.6} [Avena sativa=oats, Garry, Peptide Partial, 24 aa] gi|619331|gb|AAB31385.1||bbm|346701|bbs|152843[619331]

236: <u>1516349B</u>

BLink, Domains, Links

nm23 gene gi|226527|prf||1516349B[226527]

237: 1516349A

BLink, Domains, Links

nm23 gene gi|226526|prf||1516349A[226526]

238: CAA51527

BLink, Domains, Links

NM23H1 [Homo sapiens] gi|312824|emb|CAA51527.1|[312824]

239: AAA86745

BLink, Links

nucleoside diphosphate kinase B gi|924935|gb|AAA86745.1|[924935]

240: <u>AAA85097</u>

BLink, Domains, Links

DR-nm23 gene product gi|1051256|gb|AAA85097.1|[1051256]

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Menin, the multiple endocrine neoplasia type 1 gene product, exhibits GTP-hydrolyzing activity in the presence of the tumor metastasis

suppressor nm23.

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JOURNAL: Journal of Biological Chemistry 277 (41): p38197-38204 October

11, 2002 2002 MEDIUM: print ISSN: 0021-9258

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: MEN1, the gene responsible for multiple endocrine neoplasia type 1, is a tumor suppressor gene that encodes a *protein* called menin, of unknown function with no homology to any known *protein*. Here we demonstrate that menin interacts with a putative tumor metastasis suppressor *nm23H1*/nucleoside diphosphate (NDP) kinase A in mammalian cells. Given the roles of *nm23* as a multi-functional *protein*, we searched for the possible function of menin. Menin has no effect on the known activities of *nm23*; that is, nucleoside diphosphate kinase, *protein* kinase, or GTPase-activating *protein* for Ras-related GTPase *Rad*. However, we found that menin hydrolyzes GTP to GDP efficiently in the presence of *nm23*, whereas *nm23* or menin alone shows little or no detectable GTPase activity. Furthermore, menin contains sequence motifs similar to those found in all known GTPases or GTP-binding *proteins* and shows low affinity but specific binding to GTP/GDP. These results suggest that menin is an atypical GTPase stimulated by *nm23*.

3/3,K/2 (Item 2 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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0013093839 BIOSIS NO.: 200100265678

Tumor metastasis suppressor nm23H1 regulates Rac1 GTPase by interaction with Tiam1

AUTHOR: Otsuki Yoshiro; Tanaka Masamitsu; Yoshii Shigeto; Kawazoe Nobuko; Nakaya Kazuyasu; Sugimura Haruhiko (Reprint)

AUTHOR ADDRESS: First Department of Pathology, Hamamatsu University School of Medicine, 3600 Handa-cho, Hamamatsu, 431-3192, Japan**Japan JOURNAL: Proceedings of the National Academy of Sciences of the United States of America 98 (8): p4385-4390 April 10, 2001 2001

MEDIUM: print ISSN: 0027-8424

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The putative tumor metastasis suppressor *nm23H1* was originally identified in murine melanomas by subtraction cloning. It displays nucleoside diphosphate kinase activity and regulates cellular events, including growth and development. Recently *nm23H1* has been reported to also act as a GTPase-activating *protein* of the Ras-related GTPase *Rad* . We attempted to determine whether *nm23H1* also regulates Rho-family GTPases. Although we were unable to detect a direct association between *nm23H1* and Rho-family GTPases, *nm23H1* was shown to be associated with a Racl-specific nucleotide exchange factor, Tiaml, by interaction with its amino-terminal region in extracts from the cells expressing exogenous Tiaml and from native tissue. Overexpression of *nm23H1* inhibited the Tiam1-induced production of GTP-bound Racl and activation of c-Jun kinase. On the other hand, forced overexpression of the wild type, but not the kinase-inactivated mutant of *nm23H1*, converted the GDP-bound forms of Racl, Cdc42, and RhoA to their GTP-bound forms in vitro by its nucleoside diphosphate kinase activity, but *nm23H1* alone apparently did not produce the GTP-bound form of these GTPases in vivo. These results

suggest that *nm23H1* negatively regulates Tiam1 and inhibits Rac1 activation in vivo. Moreover, adhesion-stimulated membrane ruffles of Rat1 fibroblasts were reduced by overexpression of *nm23H1*. Based on these observations, we concluded that we had identified a function of *nm23H1* as a regulator of Rac1 and that it may be related to the effect of *nm23H1* as a tumor metastasis suppressor.

3/3,K/3 (Item 3 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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0013012947 BIOSIS NO.: 200100184786

Regulation of growth and tumorigenicity of breast cancer cells by the low molecular weight GTPase Rad and Nm23

AUTHOR: Tseng Yu-Hua; Vicent David; Zhu Jianhua; Niu Yulian; Adeyinka Adewale; Moyers Julie S; Watson Peter H; Kahn C Ronald (Reprint) AUTHOR ADDRESS: Joslin Diabetes Center, One Joslin Place, Boston, MA, 02215, USA**USA

JOURNAL: Cancer Research 61 (5): p2071-2079 March 1, 2001 2001

MEDIUM: print ISSN: 0008-5472

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: *Rad* is the prototypic member of a family of novel Ras-related GTPases that is normally expressed in heart, skeletal muscle, and lung and that has been shown to exhibit a novel form of bi-directional interaction with the *nm23* metastasis suppressor. In the present study, we have investigated the expression of *Rad* in normal and neoplastic breast tissues by Western blot and immunohistochemistry and the functional effect of altered *Rad* expression in breast cancer cell lines. We found that, although *Rad* is frequently expressed in normal breast tissue (23/30 *Rad*+ve), expression is usually lost in adjacent invasive carcinoma (8/30 *Rad*+ve; P < 0.0001). However, where *Rad* expression persists in a small proportion of tumors, it is associated with higher grade, larger size, and extensive axillary nodal involvement (n = 48; P = 0.035, P = 0.016, P = 0.022, respectively). Furthermore,*Rad* is also highly expressed in a breast cancer cell line with high tumorigenic and metastatic potential (MDA-MB231). To further examine the role of *Rad* in breast cancer, we stably transfected a *Rad*-ve breast cancer cell line (MDA-MB435). We observed an increase in growth and marked increased colony formation in soft agar in vitro (P < 0.05) and an increase in tumor growth rate in nude mice (P < 0.05). Moreover, coexpression of *nm23* with wild-type *Rad* inhibited the effect of *Rad* on growth of these cells in culture and markedly inhibited tumor growth in vivo. Additional transfection studies with mutated *Rad* cDNAs revealed that the growth-promoting effects of *Rad* appeared to be mediated through its NH2- and COOH-terminal regions, rather than its GTPase domain, and might involve acceleration of cell cycle transition. These findings suggest that *Rad* may act as an oncogenic *protein* in breast tissues and demonstrate a potential mechanism by which interaction between *Rad* and *nm23* may regulate growth and tumorigenicity of breast cancer.

3/3,K/4 (Item 4 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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0012369749 BIOSIS NO.: 200000088062

Interaction of the Ras-related *protein* associated with diabetes *Rad* and the putative tumor metastasis suppressor *NM23* provides a novel mechanism of GTPase regulation

AUTHOR: Zhu Jianhua; Tseng Yu-Hua; Kantor Jason D; Rhodes Christopher J; Zetter Bruce R; Moyers Julie S; Kahn C Ronald (Reprint)

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JOURNAL: Proceedings of the National Academy of Sciences of the United

States of America 96 (26): p14911-14918 Dec. 21, 1999 1999

MEDIUM: print ISSN: 0027-8424

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

Interaction of the Ras-related *protein* associated with diabetes *Rad* and the putative tumor metastasis suppressor *NM23* provides a novel mechanism of GTPase regulation

ABSTRACT: *Rad* is the prototypic member of a new class of Ras-related GTPases. Purification of the GTPase-activating *protein* (GAP) for *Rad* revealed *nm23*, a putative tumor metastasis suppressor and a development gene in Drosophila. Antibodies against *nm23* depleted *Rad*-GAP activity from human skeletal muscle cytosol, and bacterially expressed *nm23* reconstituted the activity. The GAP activity of *nm23* was specific for *Rad*, was absent with the S105N putative dominant negative mutant of *Rad*, and was reduced with mutations of *nm23*. In the presence of ATP, GDPcntdotRad was also reconverted to GTPcntdotRad by the nucleoside diphosphate (NDP) kinase activity of *nm23*. Simultaneously, *Rad* regulated *nm23* by enhancing its NDP kinase activity and decreasing its autophosphorylation. Melanoma cells transfected with wild-type *Rad*, but not the S105N-*Rad*, showed enhanced DNA synthesis in response to serum; this effect was lost with coexpression of *nm23*. Thus, the interaction of *nm23* and *Rad* provides a potential novel mechanism for bidirectional, bimolecular regulation in which *nm23* stimulates both GTP hydrolysis and GTP loading of *Rad* whereas *Rad* regulates activity of *nm23*. This interaction may play important roles in the effects of *Rad* on glucose metabolism and the effects of *nm23* on tumor metastasis and developmental regulation.

3/3,K/5 (Item 5 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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0011609263 BIOSIS NO.: 199800403510

Effects of phosphorylation on function of the Rad GTPase

AUTHOR: Moyers Julie S; Zhu Jianhua; Kahn C Ronald (Reprint)

AUTHOR ADDRESS: Res. Div., Joslin Diabetes Center, One Joslin Place,

Boston, MA 02215, USA**USA

JOURNAL: Biochemical Journal 333 (3): p609-614 Aug. 1, 1998 1998

MEDIUM: print ISSN: 0264-6021

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: *Rad*, Gem and Kir possess unique structural features in comparison with other Ras-like GTPases, including a C-terminal 31-residue extension that lacks typical prenylation motifs. We have recently shown that *Rad* and Gem bind calmodulin in a Ca2+-dependent manner via this C-terminal extension, involving residues 278-297 in human *Rad*. This domain also contains several consensus sites for serine phosphorylation, and *Rad* is complexed with calmodulin-dependent *protein* kinase II (CaMKII) in C2C12 cells. Here we show that *Rad* serves as a substrate for phosphorylation by CaMKII, cAMP-dependent *protein* kinase (PKA), *protein* kinase C (PKC) and casein kinase II (CKII) with stoichiometries in vitro of 0.2-1.3 mol of phosphate/mol of *Rad*. By deletion and point mutation analysis we show that phosphorylation by CaMKII and PKA occurs on a single serine residue at position 273, whereas PKC and CKII phosphorylate multiple C-terminal serine residues, including Ser214, Ser257, Ser273, Ser290 and Ser299. Incubation of *Rad* with PKA decreases

GTP binding by 60-70%, but this effect seems to be independent of phosphorylation, as it is observed with the Ser273 fwdarw Ala mutant of *Rad* containing a mutation at the site of PKA phosphorylation. The remainder of the serine kinases have no effect on *Rad* GTP binding, intrinsic GTP hydrolysis or GTP hydrolysis stimulated by the putative tumour metastasis suppressor *nm23*. However, phosphorylation of *Rad* by PKC and CKII abolishes the interaction of *Rad* with calmodulin. These findings suggest that the binding of *Rad* to calmodulin, as well as its ability to bind GTP, might be regulated by the activation of several serine kinases.

3/3,K/6 (Item 1 from file: 357)
DIALOG(R)File 357:Derwent Biotech Res.
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0230112 DBR Accession No.: 99-00213 PATENT

New method of modulating the activity of nm23 in individuals at risk from proliferative disorders - Rad mutant protein, antisense nucleic acid and antibody, used for diabetes, obesity, wound healing, tissue replacement and cancer therapy, gene therapy and drug screening

AUTHOR: Kahn C R; Zhu J

CORPORATE SOURCE: Boston, MA, USA.

PATENT ASSIGNEE: Joslin-Diabetes-Center 1998

PATENT NUMBER: WO 9844088 PATENT DATE: 981008 WPI ACCESSION NO.:

98-542695 (9846)

PRIORITY APPLIC. NO.: US 43983 APPLIC. DATE: 970403 NATIONAL APPLIC. NO.: WO 98US6521 APPLIC. DATE: 980402

LANGUAGE: English

ABSTRACT: A new method of modulating the activity of *nm23* in individuals at risk from proliferative disorders involves modulating the level of *Rad* activity. Activity of *nm23* in a cell or subject may be increased by administering a *Rad* *protein* or DNA encoding a *Rad* *protein*, and the level can be decreased by adding antisense nucleic acid which decreases *Rad* expression, by administering an anti-*Rad* antibody or a DNA sequence encoding the antibody, or by administering a *Rad* mutant or DNA encoding a *Rad* mutant. The new method may be used to treat diabetes, obesity, neuron or muscle cell development disorders, wound healing and tissue replacement, or various cancers...

... whether a subject is at risk of developing e.g. cancer, or for evaluating the ability of a test compound to modulate the interaction between *Rad* and *nm23* *proteins*. (32pp)

DESCRIPTORS: human *nm23*, *Rad* *protein* interaction modulation, *Rad* mutant *protein*, antisense nucleic acid, antibody, appl. diabetes, obesity, neuron, muscle cell development disorder, wound healing, tissue replacement, cancer susceptibility det., therapy, gene therapy, drug screening vulnerary...

3/3,K/7 (Item 1 from file: 399)
DIALOG(R)File 399:CA SEARCH(R)

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129270613 CA: 129(21)270613v PATENT

Modulating the Rad-nm23 interaction in subjects at risk for proliferative disorders

INVENTOR (AUTHOR): Kahn, C. Ronald; Zhu, Jinhua

LOCATION: USA

ASSIGNEE: Joslin Diabetes Center, Inc.

PATENT: PCT International; WO 9844088 A2 DATE: 19981008 APPLICATION: WO 98US6521 (19980402) *US 43983 (19970403)

PAGES: 32 pp. CODEN: PIXXD2 LANGUAGE: English CLASS: C12N-000/A

DESIGNATED COUNTRIES: AL; AM; AT; AU; AZ; BA; BB; BG; BR; BY; CA; CH; CN; CZ; DE; DK; EE; ES; FI; GB; GE; GH; GM; GW; HU; ID; IL; IS; JP; KE; KG; KP; KR; KZ; LC; LK; LR; LS; LT; LU; LV; MD; MG; MK; MN; MW; MX; NO; NZ; PL; PT;

RO; RU; SD; SE; SG; SI; SK; SL; TJ; TM; TR; TT; UA; UG; US; UZ; VN; YU; ZW; AM; AZ; BY; KG; KZ; MD; RU; TJ; TM DESIGNATED REGIONAL: GH; GM; KE; LS; MW; SD; SZ; UG; ZW; AT; BE; CH; CY; DE; DK; ES; FI; FR; GB; GR; IE; IT; LU; MC; NL; PT; SE; BF; BJ; CF; CG; CI; CM; GA; GN; ML; MR; NE; SN; TD; TG

3/3,K/8 (Item 1 from file: 149)
DIALOG(R)File 149:TGG Health&Wellness DB(SM)
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O1312500 SUPPLIER NUMBER: 11576824 (USE FORMAT 7 OR 9 FOR FULL TEXT)

Activation of a small GTP-binding protein by nucleoside diphosphate kinase.

Randazzo, Paul A.; Northup, John K.; Kahn, Richard A.

Science, v254, n5033, p850(4)

Nov 8,

1991

PUBLICATION FORMAT: Magazine/Journal ISSN: 0036-8075 LANGUAGE: English RECORD TYPE: Fulltext TARGET AUDIENCE: Academic WORD COUNT: 2468 LINE COUNT: 00236

... Contamination of ARF, G protein, and [beta] [gamma] preparations is common.

[10] P. A. Randazzo and R. A. Kahn, unpublished data.

[11] To prepare recombinant *nm23*-H1, *nm23*-H2, and *nm23*-1, the coding regions were amplified by the polymerase chain reaction (PCR) with cDNAs (supplied by P. Steeg) as template and synthetic oligonucleotides that incorporate...

...fragments were inserted into pET3C (supplied by W. Studier) at Nde I and Bam HI sites. BL21 (DE3) cells, were transfected with the plasmid and *protein* expression was induced with IPTG. The bacteria were lysed by sonication and after centrifugation (45 min at 100,000g), ammonium sulfate (40%) was added to...

...was suspended in and dialyzed against 20 mM tris (pH 7.4) containing NaCl (0.9%). The dialysate was fractionated on hydroxylapatite (5 ml, Bio-*Rad* HTP), equilibrated in a buffer (TED) containing 20 mM tris (pH 7.4), 1 mM EDTA, 1 mM DTT. The column was washed with TED4.3, and 23 mg of *nm23*-H1, *nm23*-H2, and *nm23*-1, respectively, consistent with their level of expression in BL21 (DE3) cells. Specific activities determined with deoxythymidine 5'-diphosphate as a substrate and the coupled...

...Parks, Methods Enzymol. 51, 376 (1978)] were 482 U/mg, 740 U/mg, 724 U/mg, and 257 U/mg at 20[degrees]C for *nm23*-H1, *nm23*-H2, *nm23*-1, and Sigma enzyme N-2635 (bovine liver), respectively. The *proteins* were estimated to be 90% pure by staining with Coomassie blue.

[12] To determine kinetic parameters, [[alpha].-sup.32]P]GDP or [[alpha].-sup.32...

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?rd
...completed examining records
              10 RD (unique items)
      S5
?s s5 not s3
                 S5
              10
                  $3
               8
                 S5 NOT S3
      S6
?show rilds;ds;t/3,k/all
>>>Invalid SHOW option: RILDS
                Description
        Items
Set
                NM23? (S) (PROTEIN? OR POLYPEPTIDE? OR PEPTIDE?)
         4029
S1
                S1 (S) (RAD OR (RAS RELATED PROTEIN ASSOCIATED WITH DIABET-
S2
           35
             ES))
                RD (unique items)
S3
            8
                S1 AND (RAD OR (RAS RELATED PROTEIN ASSOCIATED WITH DIABET-
S4
             ES))
S5
                RD (unique items)
           10
                S5 NOT S3
S6
>>>KWIC option is not available in file(s): 399
 6/3, K/1
             (Item 1 from file: 5)
DIALOG(R)File
                5:Biosis Previews(R)
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             BIOSIS NO.: 200200412105
0013818594
Integrin cytoplasmic domain-associated *protein* lalpha (ICAP-lalpha)
  interacts directly with the metastasis suppressor *nm23*-H2, and both
  *proteins* are targeted to newly formed cell adhesion sites upon integrin
  engagement
AUTHOR: Fournier Henri-Noel; Dupe-Manet Sandra; Bouvard Daniel; Lacombe
  Marie-Lise; Marie Christiane; Block Marc R (Reprint); Albiges-Rizo
  Corinne
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  Cellulaires, Faculte de Medecine de Grenoble, Institut Albert Bonniot,
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JOURNAL: Journal of Biological Chemistry 277 (23): p20895-20902 June 7,
2002 2002
MEDIUM: print
ISSN: 0021-9258
DOCUMENT TYPE: Article
RECORD TYPE: Abstract
LANGUAGE: English
```

Integrin cytoplasmic domain-associated *protein* lalpha (ICAP-lalpha) interacts directly with the metastasis suppressor *nm23*-H2, and both *proteins* are targeted to newly formed cell adhesion sites upon integrin engagement

ABSTRACT: Cell adhesion-dependent signaling implicates cytoplasmic *proteins* interacting with the intracellular tails of integrins. Among those, the integrin cytoplasmic domain-associated *protein* 1alpha (ICAP-lalpha) has been shown to interact specifically with the betal integrin cytoplasmic domain. Although it is likely that this *protein* plays an important role in controlling cell adhesion and migration, little is known about its actual function. To search for potential ICAP-lalpha-binding *proteins*, we used a yeast two-hybrid screen and identified the human metastatic suppressor *protein* *nm23*-H2 as a new partner of ICAP-1alpha. This direct interaction was confirmed in vitro, using purified recombinant ICAP-1alpha and *nm23*-H2, and by co-immunoprecipitation from CHO cell lysates over-expressing ICAP-lalpha. The physiological relevance of this interaction is provided by confocal fluorescence microscopy, which shows that ICAP-lalpha and *nm23*-H2 are co-localized in lamellipodia during the early stages of cell spreading. These adhesion sites are enriched in occupied betal integrins and precede the formation of focal adhesions devoid of ICAP-lalpha and *nm23*-H2, indicating the dynamic segregation of components of matrix adhesions.

This peripheral staining of ICAP-lalpha and *nm23*-H2 is only observed in cells spreading on fibronectin and collagen and is absent in cells spreading on poly-L-lysine, vitronectin, or laminin. This is consistent with the fact that targeting of both ICAP-lalpha and *nm23*-H2 to the cell periphery is dependent on betal integrin engagement rather than being a consequence of cell adhesion. This finding represents the first evidence that the tumor suppressor *nm23*-H2 could act on betal integrin-mediated cell adhesion by interacting with one of the integrin partners, ICAP-lalpha.

DESCRIPTORS:

CHEMICALS & BIOCHEMICALS: ...Bio-*Rad* Laboratories...

6/3,K/2 (Item 1 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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11570553 Genuine Article#: 668WU No. References: 83

Title: Nucleoside diphosphate kinases in mammalian signal transduction systems: Recent development and perspective

Author(s): Kimura N (REPRINT) ; Shimada N; Ishijima Y; Fukuda M; Takagi Y; Ishikawa N

Corporate Source: Tokyo Metropolitan Inst Gerontol, Cellular Signaling Res Grp, Itabashi Ku, 35-2 Sakaecho/Tokyo 1730015//Japan/ (REPRINT); Tokyo Metropolitan Inst Gerontol, Cellular Signaling Res Grp, Itabashi Ku, Tokyo 1730015//Japan/

Journal: JOURNAL OF BIOENERGETICS AND BIOMEMBRANES, 2003, V35, N1 (FEB), P 41-47

ISSN: 0145-479X Publication date: 20030200

Publisher: KLUWER ACADEMIC/PLENUM PUBL, 233 SPRING ST, NEW YORK, NY 10013 USA

Language: English Document Type: ARTICLE (ABSTRACT AVAILABLE)

...Abstract: their protein structural information and its significance was extended further on the basis of recent findings obtained with small molecular weight G proteins such as *Rad*, menin, and Rac. Meanwhile, observations suggesting involvement of NDP kinases in the regulation of cell growth and differentiation led to the realization that NDP kinases

...Identifiers--NERVE GROWTH-FACTOR; METASTASIS SUPPRESSOR *NM23*; PC12
PHEOCHROMOCYTOMA CELLS; ENDOCRINE NEOPLASIA TYPE-1; GTP-BINDING
PROTEIN(GS); TUMOR-METASTASIS; NEURONAL DIFFERENTIATION;
CRYSTAL-STRUCTURE; ADENYLATE-CYCLASE; ADP-RIBOSYLATION